

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/562,601

Source:

1 FWD

Date Processed by STIC:

3/5/07

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

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1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
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Revised 01/10/06



IFWO

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/562,601

DATE: 03/05/2007
TIME: 13:44:39

Input Set : N:\efs\03_05_07\10562601_efs\sequencelisting.txt
Output Set: N:\CRF4\03052007\J562601.raw

3 <110> APPLICANT: Hengst den, Christiaan D.
4 Gajic, Olivera
5 Kuipers, Oscar P.
6 Kok, Jan
7 Sikkema, Jan
8 Geurts, Johannes M.W.
9 Nauta, Arjen
11 <120> TITLE OF INVENTION: Methods and means for regulating gene expression
13 <130> FILE REFERENCE: P63590US00
15 <140> CURRENT APPLICATION NUMBER: US 10/562,601
16 <141> CURRENT FILING DATE: 2005-12-28
18 <150> PRIOR APPLICATION NUMBER: PCT/NL2004/000474
19 <151> PRIOR FILING DATE: 2004-07-02
21 <150> PRIOR APPLICATION NUMBER: EP 03077074.7
22 <151> PRIOR FILING DATE: 2003-07-02
24 <160> NUMBER OF SEQ ID NOS: 234
27 <170> SOFTWARE: PatentIn Ver. 3.3

Does Not Comply
Corrected Diskette Needed

see p2,4

ERRORED SEQUENCES

2299 <210> SEQ ID NO: 179
2300 <211> LENGTH: 26
2301 <212> TYPE: DNA
2302 <213> ORGANISM: Artificial Sequence
2304 <220> FEATURE:
2305 <223> OTHER INFORMATION: presence of the putative CodY box in arcA
E--> 2307 <400> SEQUENCE: 179 *see p2*
2308 atatttttg aaaaattttt taaaaaa

26

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2

<210> 179
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> presence of the putative CodY box in arcA

~~e~~ <400> 179
atatttttg aaaaattttt taaaaaa

26

delete extra opening bracket

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/562,601

DATE: 03/05/2007

TIME: 13:44:40

Input Set : N:\efs\03_05_07\10562601_efs\sequencelisting.txt
Output Set: N:\CRF4\03052007\J562601.raw

L:2307 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:179 differs:178
L:2739 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2732
L:2744 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2737
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2742
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2747
L:2914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:227 after pos.:0
L:2931 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:228 after pos.:0
L:2960 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:230 after pos.:0

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<210> 15
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> consensus sequence GTP binding motif in CodY homologs

<220>
<221> Variant
<222> (2)..(2)
<223> Arg

if the amino acid at this location can represent
another amino acid, use "Xaa" instead, and
explain in <220>-<223> section

<220>
<221> Variant
<222> (3)..(3) same
<223> Met, Gly, Ile, Lys, Gln

<220>
<221> Variant
<222> (4)..(4) same
<223> Gly

<220>
<221> Variant
<222> (5)..(5) same
<223> Thr

<220>
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<222> (7)..(7) same
<223> Ser

<400> 15
Gly Gly Glu Arg Leu Gly Thr Thr
1 5